

SEQUENCE LISTING

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Bartel, Paul L.

<120> Protein-Protein Interactions

<130> 2318-290-II

<150> US 60/259,572

<151> 2001-01-04

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<170> PatentIn version 3.0

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120

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172

Met Ala Ser Thr Asn Ala Glu
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220

Ser Gln Leu Gln Arg Ile Ile Arg Asp Leu Gln Asp Ala Val Thr Glu

10

15

20

cta agc aaa gaa ttt cag gaa gca ggg gaa ccc atc acg gat gac agc

268

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30

35

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 Asp Gln Lys Glu Lys Ala Thr Leu Leu Gly Asn Lys Lys Asp Tyr Trp
 60 65 70

gat tac ttc tgt gcc tgc ctg gcc aag gtg aaa gga gcc aat gat ggg 412
 Asp Tyr Phe Cys Ala Cys Leu Ala Lys Val Lys Gly Ala Asn Asp Gly
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 Ile Arg Phe Val Lys Ser Ile Ser Glu Leu Arg Thr Ser Leu Gly Lys
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gga aga gca ttt att cgc tac tcc ttg gtg cac cag agg ttg gca gac 508
 Gly Arg Ala Phe Ile Arg Tyr Ser Leu Val His Gln Arg Leu Ala Asp
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 Thr Leu Gln Gln Cys Phe Met Asn Thr Lys Val Thr Ser Asp Trp Tyr
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 Tyr Ala Arg Ser Pro Phe Leu Gln Pro Lys Leu Ser Ser Asp Ile Val
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 Gly Gln Leu Tyr Glu Leu Thr Glu Val Gln Phe Asp Leu Ala Ser Arg
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 Gly Phe Asp Leu Asp Ala Ala Trp Pro Thr Phe Ala Arg Arg Thr Leu
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 Thr Thr Gly Ser Ser Ala Tyr Leu Trp Lys Pro Pro Ser Arg Ser Ser
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agc atg agc agc ttg gtg agc agc tac ctg cag act caa gag atg gtg 796
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 Ser Asn Phe Asp Leu Asn Ser Pro Leu Asn Asn Glu Ala Leu Glu Gly
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ttt gat gag atg cga cta gag ctg gac cag ttg gag gtg cgg gag aag 892
 Phe Asp Glu Met Arg Leu Glu Leu Asp Gln Leu Glu Val Arg Glu Lys
 235 240 245

cag cta cgg gag cgc atg cag cag ctg gac aga gag aac cag gag ctg 940
 Gln Leu Arg Glu Arg Met Gln Gln Leu Asp Arg Glu Asn Gln Glu Leu
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agg gca gct gtc agc cag caa ggg gag caa ctg cag aca gag agg gag 988
 Arg Ala Ala Val Ser Gln Gln Gly Glu Gln Leu Gln Thr Glu Arg Glu
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 Arg Gly Arg Thr Ala Ala Glu Asp Asn Val Arg Leu Thr Cys Leu Val
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gct gag ctc cag aag cag tgg gag gtc acc cag gcc acc cag aac act 1084
Ala Glu Leu Gln Lys Gln Trp Glu Val Thr Gln Ala Thr Gln Asn Thr
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Glu Lys Glu Glu Asp Tyr His Thr Ala Leu Arg Arg Leu Glu Ser Met
330 335 340

ctg cag ccc ttg gca cag gag ctt gag gcc aca cgg gac tca ctg gac 1228
Leu Gln Pro Leu Ala Gln Glu Leu Glu Ala Thr Arg Asp Ser Leu Asp
345 350 355

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Lys Lys Asn Gln His Leu Ala Ser Phe Pro Gly Trp Leu Ala Met Ala
360 365 370 375

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Gln Gln Lys Ala Asp Thr Ala Ser Asp Thr Lys Gly Arg Gln Glu Pro
380 385 390

att ccc agt gat gcg gcc cag gag atg cag gag cta ggg gag aag ctt 1372
Ile Pro Ser Asp Ala Ala Gln Glu Met Gln Glu Leu Gly Glu Lys Leu
395 400 405

caa gcc cta gaa agg gag aga acc aag gtc gag gag gtc aac aga cag 1420
Gln Ala Leu Glu Arg Glu Arg Thr Lys Val Glu Glu Val Asn Arg Gln
410 415 420

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Gln Ser Ala Gln Leu Glu Gln Leu Val Lys Glu Leu Gln Leu Lys Glu
425 430 435

gat gcc cgg gcc agc ctg gag cgc ctg gtg aag gag atg gcc cca ctc 1516
Asp Ala Arg Ala Ser Leu Glu Arg Leu Val Lys Glu Met Ala Pro Leu
440 445 450 455

cag gag gag ttg tct ggg aag gga cag gag gca gac cag ctc tgg cga 1564
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204070 "EHEE001"

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